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<110> CONNAUGHT LABORATORIES LIMITED
LOOSMORE, Sheena M.
SASAKI, Ken
YANG, Yan-Ping
KLEIN, Michel H.

<120> RECOMBINANT HIGH MOLECULAR WEIGHT MAJOR OUTER MEMBRANE
PROTEIN OF MORAXELLA

<130> 1038-1063 MIS

<140> PCT/CA00/00870

<141> 2000-07-26

<150> 09/361,619

<151> 1999-07-27

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<170> PatentIn Ver. 2.1

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gaagtaacga	acttggttgg	tcttggtaat	gctggtaagt	ataacgctga	cggcaatcag	4740
gtaaacattg	ccgacatcaa	aaaagaccca	aattcagggt	catcatctaa	ccgcactgtc	4800
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ggtggtatatac	aagtgggctg	ggataaagac	ggcaacgcta	acggcgattt	aagcaatggt	4920
tgggtcaaaa	cccaaaaaga	tggcagcaaa	aaagccctgc	tcgccactta	taacgccgca	4980
ggtcagacca	actatttgac	caacaacccc	gcagaagcca	ttgacagaat	aaatgaacaa	5040
ggtatccgct	tcttccatgt	caacgatggc	aatcaagagc	ctgtggtaca	agggcgtaac	5100
ggcattgact	caagtgcctc	aggcaagcac	tcagtggcga	taggtttcca	ggccaaggca	5160
gatggtgaag	ccgccgttgc	cataggcaga	caaaccacag	caggcaacca	atccatcgcc	5220
atcggtgata	acgcacaagc	cacgggcgat	caatccatcg	ccatcggtac	aggcaatgtg	5280
gtagcaggta	agcactctgg	tgccatcggc	gacccaagca	ctgttaaggc	tgataacagt	5340
tacagtgtgg	gtaataacaa	caggtttacc	cgatgccactc	aaaccgatgt	ctttgggtgtg	5400
ggcaataaca	tcaccgtgac	cgaaagtaac	tcggttgctc	taggttcaaa	ctctgccatc	5460
agtgcaggca	cacacgcagg	cacacaagcc	aaaaaatctg	acggcacagc	aggtacaacc	5520
accacagcag	gtgcaaccgg	tacggttaaa	ggctttgctg	gacaaaacggc	ggttgggtgcg	5580
gtctccgtgg	gtgcctcagg	tgctgaacgc	cgtatccaaa	atgtggcagc	aggtgaggtc	5640
agtggccacca	gcaccgatgc	ggtcaatggt	agccagttgt	acaaagccac	ccaaagcatt	5700
gccaaacgcaa	ccaatgagct	tgaccatcgt	atccacaaa	acgaaaataa	ggccaatgca	5760
gggatttcac	cagcgatggc	gatggcgctc	atgccacaag	cctacattcc	tggcagatcc	5820
atggttaccg	ggggtattgc	caccacaaac	ggtcaagggtg	cggtggcagt	gggactgtcg	5880

Substitute SeqListing.txt

aagctgtcgg ataatgggtca atgggtatttt aaaatcaatg gttcagccga tacccaaggc 5940
catgtagggg cggcagttgg tgcaggtttt cacttt 5976

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<212> PRT
<213> Moraxella catarrhalis

<400> 3
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Thr Lys His Ile Ala Ile Gly Glu Gln Asn Gln Pro Arg Arg Ser Gly
20 25 30
Thr Ala Lys Ala Asp Gly Asp Arg Ala Ile Ala Ile Gly Glu Asn Ala
35 40 45
Asn Ala Gln Gly Gly Gln Ala Ile Ala Ile Gly Ser Ser Asn Lys Thr
50 55 60
Val Asn Gly Ser Ser Leu Asp Lys Ile Gly Thr Asp Ala Thr Gly Gln
65 70 75 80
Glu Ser Ile Ala Ile Gly Gly Asp Val Lys Ala Ser Gly Asp Ala Ser
85 90 95
Ile Ala Ile Gly Ser Asp Asp Leu His Leu Leu Asp Gln His Gly Asn
100 105 110
Pro Lys His Pro Lys Gly Thr Leu Ile Asn Asp Leu Ile Asn Gly His
115 120 125
Ala Val Leu Lys Glu Ile Arg Ser Ser Lys Asp Asn Asp Val Lys Tyr
130 135 140
Arg Arg Thr Thr Ala Ser Gly His Ala Ser Thr Ala Val Gly Ala Met
145 150 155 160
Ser Tyr Ala Gln Gly His Phe Ser Asn Ala Phe Gly Thr Arg Ala Thr
165 170 175
Ala Lys Ser Ala Tyr Ser Leu Ala Val Gly Leu Ala Ala Thr Ala Glu
180 185 190
Gly Gln Ser Thr Ile Ala Ile Gly Ser Asp Ala Thr Ser Ser Ser Leu
195 200 205
Gly Ala Ile Ala Leu Gly Ala Gly Thr Arg Ala Gln Leu Gln Gly Ser
210 215 220
Ile Ala Leu Gly Gln Gly Ser Val Val Thr Gln Ser Asp Asn Asn Ser
225 230 235 240
Arg Pro Ala Tyr Thr Pro Asn Thr Gln Ala Leu Asp Pro Lys Phe Gln
245 250 255
Ala Thr Asn Asn Thr Lys Ala Gly Pro Leu Ser Ile Gly Ser Asn Ser
260 265 270
Ile Lys Arg Lys Ile Ile Asn Val Gly Ala Gly Val Asn Lys Thr Asp
275 280 285

Substitute SeqListing.txt

Ala Val Asn Val Ala Gln Leu Glu Ala Val Val Lys Trp Ala Lys Glu
290 295 300

Arg Arg Ile Thr Phe Gln Gly Asp Asp Asn Ser Thr Asp Val Lys Ile
305 310 315 320

Gly Leu Asp Asn Thr Leu Thr Ile Lys Gly Gly Ala Glu Thr Asn Ala
325 330 335

Leu Thr Asp Asn Asn Ile Gly Val Val Lys Glu Ala Asp Asn Ser Gly
340 345 350

Leu Lys Val Lys Leu Ala Lys Thr Leu Asn Asn Leu Thr Glu Val Asn
355 360 365

Thr Thr Thr Leu Asn Ala Thr Thr Thr Val Lys Val Gly Ser Ser Ser
370 375 380

Ser Thr Thr Ala Glu Leu Leu Ser Asp Ser Leu Thr Phe Thr Gln Pro
385 390 395 400

Asn Thr Gly Ser Gln Ser Thr Ser Lys Thr Val Tyr Gly Val Asn Gly
405 410 415

Val Lys Phe Thr Asn Asn Ala Glu Thr Thr Ala Ala Ile Gly Thr Thr
420 425 430

Arg Ile Thr Arg Asp Lys Ile Gly Phe Ala Arg Asp Gly Asp Val Asp
435 440 445

Glu Lys Gln Ala Pro Tyr Leu Asp Lys Lys Gln Leu Lys Val Gly Ser
450 455 460

Val Ala Ile Thr Ile Asp Asn Gly Ile Asp Ala Gly Asn Lys Lys Ile
465 470 475 480

Ser Asn Leu Ala Lys Gly Ser Ser Ala Asn Asp Ala Val Thr Ile Glu
485 490 495

Gln Leu Lys Ala Ala Lys Pro Thr Leu Asn Ala Gly Ala Gly Ile Ser
500 505 510

Val Thr Pro Thr Glu Ile Ser Val Asp Ala Lys Ser Gly Asn Val Thr
515 520 525

Ala Pro Thr Tyr Asn Ile Gly Val Lys Thr Thr Glu Leu Asn Ser Asp
530 535 540

Gly Thr Ser Asp Lys Phe Ser Val Lys Gly Ser Gly Thr Asn Asn Ser
545 550 555 560

Leu Val Thr Ala Glu His Leu Ala Ser Tyr Leu Asn Glu Val Asn Arg
565 570 575

Thr Ala Asp Ser Ala Leu Gln Ser Phe Thr Val Lys Glu Glu Asp Asp
580 585 590

Asp Asp Ala Asn Ala Ile Thr Val Ala Lys Asp Thr Thr Lys Asn Ala
595 600 605

Gly Ala Val Ser Ile Leu Lys Leu Lys Gly Lys Asn Gly Leu Thr Val
610 615 620

Substitute SeqListing.txt

Ala Thr Lys Lys Asp Gly Thr Val Thr Phe Gly Leu Ser Gln Asp Ser
625 630 635 640

Gly Leu Thr Ile Gly Lys Ser Thr Leu Asn Asn Asp Gly Leu Thr Val
645 650 655

Lys Asp Thr Asn Glu Gln Ile Gln Val Gly Ala Asn Gly Ile Lys Phe
660 665 670

Thr Asn Val Asn Gly Ser Asn Pro Gly Thr Gly Ile Ala Asn Thr Ala
675 680 685

Arg Ile Thr Arg Asp Lys Ile Gly Phe Ala Gly Ser Asp Gly Ala Val
690 695 700

Asp Thr Asn Lys Pro Tyr Leu Asp Gln Asp Lys Leu Gln Val Gly Asn
705 710 715 720

Val Lys Ile Thr Asn Thr Gly Ile Asn Ala Gly Gly Lys Ala Ile Thr
725 730 735

Gly Leu Ser Pro Thr Leu Pro Ser Ile Ala Asp Gln Ser Ser Arg Asn
740 745 750

Ile Glu Leu Gly Asn Thr Ile Gln Asp Lys Asp Lys Ser Asn Ala Ala
755 760 765

Ser Ile Asn Asp Ile Leu Asn Thr Gly Phe Asn Leu Lys Asn Asn Asn
770 775 780

Asn Pro Ile Asp Phe Val Ser Thr Tyr Asp Ile Val Asp Phe Ala Asn
785 790 795 800

Gly Asn Ala Thr Thr Ala Thr Val Thr His Asp Thr Ala Asn Lys Thr
805 810 815

Ser Lys Val Val Tyr Asp Val Asn Val Asp Asp Thr Thr Ile His Leu
820 825 830

Thr Gly Thr Asp Asp Asn Lys Lys Leu Gly Val Lys Thr Thr Lys Leu
835 840 845

Asn Lys Thr Ser Ala Asn Gly Asn Thr Ala Thr Asn Phe Asn Val Asn
850 855 860

Ser Ser Asp Glu Asp Ala Leu Val Asn Ala Lys Asp Ile Ala Glu Asn
865 870 875 880

Leu Asn Thr Leu Ala Lys Glu Ile His Thr Thr Lys Gly Thr Ala Asp
885 890 895

Thr Ala Leu Gln Thr Phe Thr Val Lys Lys Val Asp Glu Asn Asn Asn
900 905 910

Ala Asp Asp Ala Asn Ala Ile Thr Val Gly Gln Lys Asn Ala Asn Asn
915 920 925

Gln Val Asn Thr Leu Thr Leu Lys Gly Glu Asn Gly Leu Asn Ile Lys
930 935 940

Thr Asp Lys Asn Gly Thr Val Thr Phe Gly Ile Asn Thr Thr Ser Gly
945 950 955 960

Substitute SeqListing.txt

Leu Lys Ala Gly Lys Ser Thr Leu Asn Asp Gly Gly Leu Ser Ile Lys
 965 970 975
 Asn Pro Thr Gly Ser Glu Gln Ile Gln Val Gly Ala Asp Gly Val Lys
 980 985 990
 Phe Ala Lys Val Asn Asn Asn Gly Val Val Gly Ala Gly Ile Asp Gly
 995 1000 1005
 Thr Thr Arg Ile Thr Arg Asp Glu Ile Gly Phe Thr Gly Thr Asn Gly
 1010 1015 1020
 Ser Leu Asp Lys Ser Lys Pro His Leu Ser Lys Asp Gly Ile Asn Ala
 1025 1030 1035 1040
 Gly Gly Lys Lys Ile Thr Asn Ile Gln Ser Gly Glu Ile Ala Gln Asn
 1045 1050 1055
 Ser His Asp Ala Val Thr Gly Gly Lys Ile Tyr Asp Leu Lys Thr Glu
 1060 1065 1070
 Leu Glu Asn Lys Ile Ser Ser Thr Ala Lys Thr Ala Gln Asn Ser Leu
 1075 1080 1085
 His Glu Phe Ser Val Ala Asp Glu Gln Gly Asn Asn Phe Thr Val Ser
 1090 1095 1100
 Asn Pro Tyr Ser Ser Tyr Asp Thr Ser Lys Thr Ser Asp Val Ile Thr
 1105 1110 1115 1120
 Phe Ala Gly Glu Asn Gly Ile Thr Thr Lys Val Asn Lys Gly Val Val
 1125 1130 1135
 Arg Val Gly Ile Asp Gln Thr Lys Gly Leu Thr Thr Pro Lys Leu Thr
 1140 1145 1150
 Val Gly Asn Asn Asn Gly Lys Gly Ile Val Ile Asp Ser Gln Asn Gly
 1155 1160 1165
 Gln Asn Thr Ile Thr Gly Leu Ser Asn Thr Leu Ala Asn Val Thr Asn
 1170 1175 1180
 Asp Lys Gly Ser Val Arg Thr Thr Glu Gln Gly Asn Ile Ile Lys Asp
 1185 1190 1195 1200
 Glu Asp Lys Thr Arg Ala Ala Ser Ile Val Asp Val Leu Ser Ala Gly
 1205 1210 1215
 Phe Asn Leu Gln Gly Asn Gly Glu Ala Val Asp Phe Val Ser Thr Tyr
 1220 1225 1230
 Asp Thr Val Asn Phe Ala Asp Gly Asn Ala Thr Thr Ala Lys Val Thr
 1235 1240 1245
 Tyr Asp Asp Thr Ser Lys Thr Ser Lys Val Val Tyr Asp Val Asn Val
 1250 1255 1260
 Asp Asp Thr Thr Ile Glu Val Lys Asp Lys Lys Leu Gly Val Lys Thr
 1265 1270 1275 1280
 Thr Thr Leu Thr Ser Thr Gly Thr Gly Ala Asn Lys Phe Ala Leu Ser
 1285 1290 1295

Substitute SeqListing.txt

Asn Gln Ala Thr Gly Asp Ala Leu Val Lys Ala Ser Asp Ile Val Ala
 1300 1305 1310
 His Leu Asn Thr Leu Ser Gly Asp Ile Gln Thr Ala Lys Gly Ala Ser
 1315 1320 1325
 Gln Ala Asn Asn Ser Ala Gly Tyr Val Asp Ala Asp Gly Asn Lys Val
 1330 1335 1340
 Ile Tyr Asp Ser Thr Asp Asn Lys Tyr Tyr Gln Ala Lys Asn Asp Gly
 1345 1350 1355 1360
 Thr Val Asp Lys Thr Lys Glu Val Ala Lys Asp Lys Leu Val Ala Gln
 1365 1370 1375
 Ala Gln Thr Pro Asp Gly Thr Leu Ala Gln Met Asn Val Lys Ser Val
 1380 1385 1390
 Ile Asn Lys Glu Gln Val Asn Asp Ala Asn Lys Lys Gln Gly Ile Asn
 1395 1400 1405
 Glu Asp Asn Ala Phe Val Lys Gly Leu Glu Lys Ala Ala Ser Asp Asn
 1410 1415 1420
 Lys Thr Lys Asn Ala Ala Val Thr Val Gly Asp Leu Asn Ala Val Ala
 1425 1430 1435 1440
 Gln Thr Pro Leu Thr Phe Ala Gly Asp Thr Gly Thr Thr Ala Lys Lys
 1445 1450 1455
 Leu Gly Glu Thr Leu Thr Ile Lys Gly Gly Gln Thr Asp Thr Asn Lys
 1460 1465 1470
 Leu Thr Asp Asn Asn Ile Gly Val Val Ala Gly Thr Asp Gly Phe Thr
 1475 1480 1485
 Val Lys Leu Ala Lys Asp Leu Thr Asn Leu Asn Ser Val Asn Ala Gly
 1490 1495 1500
 Gly Thr Lys Ile Asp Asp Lys Gly Val Ser Phe Val Asp Ser Ser Gly
 1505 1510 1515 1520
 Gln Ala Lys Ala Asn Thr Pro Val Leu Ser Ala Asn Gly Leu Asp Leu
 1525 1530 1535
 Gly Gly Lys Val Ile Ser Asn Val Gly Lys Gly Thr Lys Asp Thr Asp
 1540 1545 1550
 Ala Ala Asn Val Gln Gln Leu Asn Glu Val Arg Asn Leu Leu Gly Leu
 1555 1560 1565
 Gly Asn Ala Gly Asn Asp Asn Ala Asp Gly Asn Gln Val Asn Ile Ala
 1570 1575 1580
 Asp Ile Lys Lys Asp Pro Asn Ser Gly Ser Ser Ser Asn Arg Thr Val
 1585 1590 1595 1600
 Ile Lys Ala Gly Thr Val Leu Gly Gly Lys Gly Asn Asn Asp Thr Glu
 1605 1610 1615
 Lys Leu Ala Thr Gly Gly Ile Gln Val Gly Val Asp Lys Asp Gly Asn
 1620 1625 1630

Substitute SeqListing.txt

Ala Asn Gly Asp Leu Ser Asn Val Trp Val Lys Thr Gln Lys Asp Gly
1635 1640 1645

Ser Lys Lys Ala Leu Leu Ala Thr Tyr Asn Ala Ala Gly Gln Thr Asn
1650 1655 1660

Tyr Leu Thr Asn Asn Pro Ala Glu Ala Ile Asp Arg Ile Asn Glu Gln
1665 1670 1675 1680

Gly Ile Arg Phe Phe His Val Asn Asp Gly Asn Gln Glu Pro Val Val
1685 1690 1695

Gln Gly Arg Asn Gly Ile Asp Ser Ser Ala Ser Gly Lys His Ser Val
1700 1705 1710

Ala Ile Gly Phe Gln Ala Lys Ala Asp Gly Glu Ala Ala Val Ala Ile
1715 1720 1725

Gly Arg Gln Thr Gln Ala Gly Asn Gln Ser Ile Ala Ile Gly Asp Asn
1730 1735 1740

Ala Gln Ala Thr Gly Asp Gln Ser Ile Ala Ile Gly Thr Gly Asn Val
1745 1750 1755 1760

Val Ala Gly Lys His Ser Gly Ala Ile Gly Asp Pro Ser Thr Val Lys
1765 1770 1775

Ala Asp Asn Ser Tyr Ser Val Gly Asn Asn Asn Gln Phe Thr Asp Ala
1780 1785 1790

Thr Gln Thr Asp Val Phe Gly Val Gly Asn Asn Ile Thr Val Thr Glu
1795 1800 1805

Ser Asn Ser Val Ala Leu Gly Ser Asn Ser Ala Ile Ser Ala Gly Thr
1810 1815 1820

His Ala Gly Thr Gln Ala Lys Lys Ser Asp Gly Thr Ala Gly Thr Thr
1825 1830 1835 1840

Thr Thr Ala Gly Ala Thr Gly Thr Val Lys Gly Phe Ala Gly Gln Thr
1845 1850 1855

Ala Val Gly Ala Val Ser Val Gly Ala Ser Gly Ala Glu Arg Arg Ile
1860 1865 1870

Gln Asn Val Ala Ala Gly Glu Val Ser Ala Thr Ser Thr Asp Ala Val
1875 1880 1885

Asn Gly Ser Gln Leu Tyr Lys Ala Thr Gln Ser Ile Ala Asn Ala Thr
1890 1895 1900

Asn Glu Leu Asp His Arg Ile His Gln Asn Glu Asn Lys Ala Asn Ala
1905 1910 1915 1920

Gly Ile Ser Ser Ala Met Ala Met Ala Ser Met Pro Gln Ala Tyr Ile
1925 1930 1935

Pro Gly Arg Ser Met Val Thr Gly Gly Ile Ala Thr His Asn Gly Gln
1940 1945 1950

Gly Ala Val Ala Val Gly Leu Ser Lys Leu Ser Asp Asn Gly Gln Trp
1955 1960 1965

Substitute SeqListing.txt

Val Phe Lys Ile Asn Gly Ser Ala Asp Thr Gln Gly His Val Gly Ala
1970 1975 1980

Ala Val Gly Ala Gly Phe His Phe
1985 1990

<210> 4

<211> 2047

<212> PRT

<213> Moraxella catarrhalis

<400> 4

Met Asn His Ile Tyr Lys Val Ile Phe Asn Lys Ala Thr Gly Thr Phe
1 5 10 15

Met Ala Val Ala Glu Tyr Ala Lys Ser His Ser Thr Gly Gly Gly Ser
20 25 30

Cys Ala Thr Gly Gln Val Gly Ser Val Cys Thr Leu Ser Phe Ala Arg
35 40 45

Ile Ala Ala Leu Ala Val Leu Val Ile Gly Ala Thr Leu Ser Gly Ser
50 55 60

Ala Tyr Ala Gln Lys Lys Asp Thr Lys His Ile Ala Ile Gly Glu Gln
65 70 75 80

Asn Gln Pro Arg Arg Ser Gly Thr Ala Lys Ala Asp Gly Asp Arg Ala
85 90 95

Ile Ala Ile Gly Glu Asn Ala Asn Ala Gln Gly Gly Gln Ala Ile Ala
100 105 110

Ile Gly Ser Ser Asn Lys Thr Val Asn Gly Ser Ser Leu Asp Lys Ile
115 120 125

Gly Thr Asp Ala Thr Gly Gln Glu Ser Ile Ala Ile Gly Gly Asp Val
130 135 140

Lys Ala Ser Gly Asp Ala Ser Ile Ala Ile Gly Ser Asp Asp Leu His
145 150 155 160

Leu Leu Asp Gln His Gly Asn Pro Lys His Pro Lys Gly Thr Leu Ile
165 170 175

Asn Asp Leu Ile Asn Gly His Ala Val Leu Lys Glu Ile Arg Ser Ser
180 185 190

Lys Asp Asn Asp Val Lys Tyr Arg Arg Thr Thr Ala Ser Gly His Ala
195 200 205

Ser Thr Ala Val Gly Ala Met Ser Tyr Ala Gln Gly His Phe Ser Asn
210 215 220

Ala Phe Gly Thr Arg Ala Thr Ala Lys Ser Ala Tyr Ser Leu Ala Val
225 230 235 240

Gly Leu Ala Ala Thr Ala Glu Gly Gln Ser Thr Ile Ala Ile Gly Ser
245 250 255

Asp Ala Thr Ser Ser Ser Leu Gly Ala Ile Ala Leu Gly Ala Gly Thr
Page 11

Substitute SeqListing.txt
265 270

260

Arg Ala Gln Leu Gln Gly Ser Ile Ala Leu Gly Gln Gly Ser Val Val
275 280 285

Thr Gln Ser Asp Asn Asn Ser Arg Pro Ala Tyr Thr Pro Asn Thr Gln
290 295 300

Ala Leu Asp Pro Lys Phe Gln Ala Thr Asn Asn Thr Lys Ala Gly Pro
305 310 315 320

Leu Ser Ile Gly Ser Asn Ser Ile Lys Arg Lys Ile Ile Asn Val Gly
325 330 335

Ala Gly Val Asn Lys Thr Asp Ala Val Asn Val Ala Gln Leu Glu Ala
340 345 350

Val Val Lys Trp Ala Lys Glu Arg Arg Ile Thr Phe Gln Gly Asp Asp
355 360 365

Asn Ser Thr Asp Val Lys Ile Gly Leu Asp Asn Thr Leu Thr Ile Lys
370 375 380

Gly Gly Ala Glu Thr Asn Ala Leu Thr Asp Asn Asn Ile Gly Val Val
385 390 395 400

Lys Glu Ala Asp Asn Ser Gly Leu Lys Val Lys Leu Ala Lys Thr Leu
405 410 415

Asn Asn Leu Thr Glu Val Asn Thr Thr Thr Leu Asn Ala Thr Thr Thr
420 425 430

Val Lys Val Gly Ser Ser Ser Ser Thr Thr Ala Glu Leu Leu Ser Asp
435 440 445

Ser Leu Thr Phe Thr Gln Pro Asn Thr Gly Ser Gln Ser Thr Ser Lys
450 455 460

Thr Val Tyr Gly Val Asn Gly Val Lys Phe Thr Asn Asn Ala Glu Thr
465 470 475 480

Thr Ala Ala Ile Gly Thr Thr Arg Ile Thr Arg Asp Lys Ile Gly Phe
485 490 495

Ala Arg Asp Gly Asp Val Asp Glu Lys Gln Ala Pro Tyr Leu Asp Lys
500 505 510

Lys Gln Leu Lys Val Gly Ser Val Ala Ile Thr Ile Asp Asn Gly Ile
515 520 525

Asp Ala Gly Asn Lys Lys Ile Ser Asn Leu Ala Lys Gly Ser Ser Ala
530 535 540

Asn Asp Ala Val Thr Ile Glu Gln Leu Lys Ala Ala Lys Pro Thr Leu
545 550 555 560

Asn Ala Gly Ala Gly Ile Ser Val Thr Pro Thr Glu Ile Ser Val Asp
565 570 575

Ala Lys Ser Gly Asn Val Thr Ala Pro Thr Tyr Asn Ile Gly Val Lys
580 585 590

Thr Thr Glu Leu Asn Ser Asp Gly Thr Ser Asp Lys Phe Ser Val Lys

Substitute SeqListing.txt
600 605

595

Gly Ser Gly Thr Asn Asn Ser Leu Val Thr Ala Glu His Leu Ala Ser
610 615 620
Tyr Leu Asn Glu Val Asn Arg Thr Ala Asp Ser Ala Leu Gln Ser Phe
625 630 635 640
Thr Val Lys Glu Glu Asp Asp Asp Asp Ala Asn Ala Ile Thr Val Ala
645 650 655
Lys Asp Thr Thr Lys Asn Ala Gly Ala Val Ser Ile Leu Lys Leu Lys
660 665 670
Gly Lys Asn Gly Leu Thr Val Ala Thr Lys Lys Asp Gly Thr Val Thr
675 680 685
Phe Gly Leu Ser Gln Asp Ser Gly Leu Thr Ile Gly Lys Ser Thr Leu
690 695 700
Asn Asn Asp Gly Leu Thr Val Lys Asp Thr Asn Glu Gln Ile Gln Val
705 710 715 720
Gly Ala Asn Gly Ile Lys Phe Thr Asn Val Asn Gly Ser Asn Pro Gly
725 730 735
Thr Gly Ile Ala Asn Thr Ala Arg Ile Thr Arg Asp Lys Ile Gly Phe
740 745 750
Ala Gly Ser Asp Gly Ala Val Asp Thr Asn Lys Pro Tyr Leu Asp Gln
755 760 765
Asp Lys Leu Gln Val Gly Asn Val Lys Ile Thr Asn Thr Gly Ile Asn
770 775 780
Ala Gly Gly Lys Ala Ile Thr Gly Leu Ser Pro Thr Leu Pro Ser Ile
785 790 795 800
Ala Asp Gln Ser Ser Arg Asn Ile Glu Leu Gly Asn Thr Ile Gln Asp
805 810 815
Lys Asp Lys Ser Asn Ala Ala Ser Ile Asn Asp Ile Leu Asn Thr Gly
820 825 830
Phe Asn Leu Lys Asn Asn Asn Asn Pro Ile Asp Phe Val Ser Thr Tyr
835 840 845
Asp Ile Val Asp Phe Ala Asn Gly Asn Ala Thr Thr Ala Thr Val Thr
850 855 860
His Asp Thr Ala Asn Lys Thr Ser Lys Val Val Tyr Asp Val Asn Val
865 870 875 880
Asp Asp Thr Thr Ile His Leu Thr Gly Thr Asp Asp Asn Lys Lys Leu
885 890 895
Gly Val Lys Thr Thr Lys Leu Asn Lys Thr Ser Ala Asn Gly Asn Thr
900 905 910
Ala Thr Asn Phe Asn Val Asn Ser Ser Asp Glu Asp Ala Leu Val Asn
915 920 925
Ala Lys Asp Ile Ala Glu Asn Leu Asn Thr Leu Ala Lys Glu Ile His

Substitute SeqListing.txt

930 935 940
 Thr Thr Lys Gly Thr Ala Asp Thr Ala Leu Gln Thr Phe Thr Val Lys
 945 950 955 960
 Lys Val Asp Glu Asn Asn Asn Ala Asp Asp Ala Asn Ala Ile Thr Val
 965 970 975
 Gly Gln Lys Asn Ala Asn Asn Gln Val Asn Thr Leu Thr Leu Lys Gly
 980 985 990
 Glu Asn Gly Leu Asn Ile Lys Thr Asp Lys Asn Gly Thr Val Thr Phe
 995 1000 1005
 Gly Ile Asn Thr Thr Ser Gly Leu Lys Ala Gly Lys Ser Thr Leu Asn
 1010 1015 1020
 Asp Gly Gly Leu Ser Ile Lys Asn Pro Thr Gly Ser Glu Gln Ile Gln
 1025 1030 1035 1040
 Val Gly Ala Asp Gly Val Lys Phe Ala Lys Val Asn Asn Asn Gly Val
 1045 1050 1055
 Val Gly Ala Gly Ile Asp Gly Thr Thr Arg Ile Thr Arg Asp Glu Ile
 1060 1065 1070
 Gly Phe Thr Gly Thr Asn Gly Ser Leu Asp Lys Ser Lys Pro His Leu
 1075 1080 1085
 Ser Lys Asp Gly Ile Asn Ala Gly Gly Lys Lys Ile Thr Asn Ile Gln
 1090 1095 1100
 Ser Gly Glu Ile Ala Gln Asn Ser His Asp Ala Val Thr Gly Gly Lys
 1105 1110 1115 1120
 Ile Tyr Asp Leu Lys Thr Glu Leu Glu Asn Lys Ile Ser Ser Thr Ala
 1125 1130 1135
 Lys Thr Ala Gln Asn Ser Leu His Glu Phe Ser Val Ala Asp Glu Gln
 1140 1145 1150
 Gly Asn Asn Phe Thr Val Ser Asn Pro Tyr Ser Ser Tyr Asp Thr Ser
 1155 1160 1165
 Lys Thr Ser Asp Val Ile Thr Phe Ala Gly Glu Asn Gly Ile Thr Thr
 1170 1175 1180
 Lys Val Asn Lys Gly Val Val Arg Val Gly Ile Asp Gln Thr Lys Gly
 1185 1190 1195 1200
 Leu Thr Thr Pro Lys Leu Thr Val Gly Asn Asn Asn Gly Lys Gly Ile
 1205 1210 1215
 Val Ile Asp Ser Gln Asn Gly Gln Asn Thr Ile Thr Gly Leu Ser Asn
 1220 1225 1230
 Thr Leu Ala Asn Val Thr Asn Asp Lys Gly Ser Val Arg Thr Thr Glu
 1235 1240 1245
 Gln Gly Asn Ile Ile Lys Asp Glu Asp Lys Thr Arg Ala Ala Ser Ile
 1250 1255 1260
 Val Asp Val Leu Ser Ala Gly Phe Asn Leu Gln Gly Asn Gly Glu Ala

Substitute SeqListing.txt

1265 1270 1275 1280
Val Asp Phe Val Ser Thr Tyr Asp Thr Val Asn Phe Ala Asp Gly Asn
1285 1290 1295
Ala Thr Thr Ala Lys Val Thr Tyr Asp Asp Thr Ser Lys Thr Ser Lys
1300 1305 1310
Val Val Tyr Asp Val Asn Val Asp Asp Thr Thr Ile Glu Val Lys Asp
1315 1320 1325
Lys Lys Leu Gly Val Lys Thr Thr Thr Leu Thr Ser Thr Gly Thr Gly
1330 1335 1340
Ala Asn Lys Phe Ala Leu Ser Asn Gln Ala Thr Gly Asp Ala Leu Val
1345 1350 1355 1360
Lys Ala Ser Asp Ile Val Ala His Leu Asn Thr Leu Ser Gly Asp Ile
1365 1370 1375
Gln Thr Ala Lys Gly Ala Ser Gln Ala Asn Asn Ser Ala Gly Tyr Val
1380 1385 1390
Asp Ala Asp Gly Asn Lys Val Ile Tyr Asp Ser Thr Asp Asn Lys Tyr
1395 1400 1405
Tyr Gln Ala Lys Asn Asp Gly Thr Val Asp Lys Thr Lys Glu Val Ala
1410 1415 1420
Lys Asp Lys Leu Val Ala Gln Ala Gln Thr Pro Asp Gly Thr Leu Ala
1425 1430 1435 1440
Gln Met Asn Val Lys Ser Val Ile Asn Lys Glu Gln Val Asn Asp Ala
1445 1450 1455
Asn Lys Lys Gln Gly Ile Asn Glu Asp Asn Ala Phe Val Lys Gly Leu
1460 1465 1470
Glu Lys Ala Ala Ser Asp Asn Lys Thr Lys Asn Ala Ala Val Thr Val
1475 1480 1485
Gly Asp Leu Asn Ala Val Ala Gln Thr Pro Leu Thr Phe Ala Gly Asp
1490 1495 1500
Thr Gly Thr Thr Ala Lys Lys Leu Gly Glu Thr Leu Thr Ile Lys Gly
1505 1510 1515 1520
Gly Gln Thr Asp Thr Asn Lys Leu Thr Asp Asn Asn Ile Gly Val Val
1525 1530 1535
Ala Gly Thr Asp Gly Phe Thr Val Lys Leu Ala Lys Asp Leu Thr Asn
1540 1545 1550
Leu Asn Ser Val Asn Ala Gly Gly Thr Lys Ile Asp Asp Lys Gly Val
1555 1560 1565
Ser Phe Val Asp Ser Ser Gly Gln Ala Lys Ala Asn Thr Pro Val Leu
1570 1575 1580
Ser Ala Asn Gly Leu Asp Leu Gly Gly Lys Val Ile Ser Asn Val Gly
1585 1590 1595 1600
Lys Gly Thr Lys Asp Thr Asp Ala Ala Asn Val Gln Gln Leu Asn Glu

Substitute SeqListing.txt

1605 1610 1615
 Val Arg Asn Leu Leu Gly Leu Gly Asn Ala Gly Asn Asp Asn Ala Asp
 1620 1625 1630
 Gly Asn Gln Val Asn Ile Ala Asp Ile Lys Lys Asp Pro Asn Ser Gly
 1635 1640 1645
 Ser Ser Ser Asn Arg Thr Val Ile Lys Ala Gly Thr Val Leu Gly Gly
 1650 1655 1660
 Lys Gly Asn Asn Asp Thr Glu Lys Leu Ala Thr Gly Gly Ile Gln Val
 1665 1670 1675 1680
 Gly Val Asp Lys Asp Gly Asn Ala Asn Gly Asp Leu Ser Asn Val Trp
 1685 1690 1695
 Val Lys Thr Gln Lys Asp Gly Ser Lys Lys Ala Leu Leu Ala Thr Tyr
 1700 1705 1710
 Asn Ala Ala Gly Gln Thr Asn Tyr Leu Thr Asn Asn Pro Ala Glu Ala
 1715 1720 1725
 Ile Asp Arg Ile Asn Glu Gln Gly Ile Arg Phe Phe His Val Asn Asp
 1730 1735 1740
 Gly Asn Gln Glu Pro Val Val Gln Gly Arg Asn Gly Ile Asp Ser Ser
 1745 1750 1755 1760
 Ala Ser Gly Lys His Ser Val Ala Ile Gly Phe Gln Ala Lys Ala Asp
 1765 1770 1775
 Gly Glu Ala Ala Val Ala Ile Gly Arg Gln Thr Gln Ala Gly Asn Gln
 1780 1785 1790
 Ser Ile Ala Ile Gly Asp Asn Ala Gln Ala Thr Gly Asp Gln Ser Ile
 1795 1800 1805
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 1810 1815 1820
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 1825 1830 1835 1840
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 1845 1850 1855
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 1875 1880 1885
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 1890 1895 1900
 Lys Gly Phe Ala Gly Gln Thr Ala Val Gly Ala Val Ser Val Gly Ala
 1905 1910 1915 1920
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 1925 1930 1935
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Substitute SeqListing.txt
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1940
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Substitute SeqListing.txt

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Substitute SeqListing.txt

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Substitute SeqListing.txt

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Substitute SeqListing.txt

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 35 40 45
 Ile Ala Ala Leu Ala Val Leu Val Ile Gly Ala Thr Leu Ser Gly Ser
 50 55 60
 Ala Tyr Ala Gln Lys Lys Asp Thr Lys His Ile Ala Ile Gly Glu Gln
 65 70 75 80
 Asn Gln Pro Arg Arg Ser Gly Thr Ala Lys Ala Asp Gly Asp Arg Ala
 85 90 95
 Ile Ala Ile Gly Glu Asn Ala Asn Ala Gln Gly Gly Gln Ala Ile Ala
 100 105 110
 Ile Gly Ser Ser Asn Lys Thr Val Asn Gly Ser Ser Leu Asp Lys Ile
 115 120 125
 Gly Thr Asp Ala Thr Gly Gln Glu Ser Ile Ala Ile Gly Gly Asp Val
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 Lys Ala Ser Gly Asp Ala Ser Ile Ala Ile Gly Ser Asp Asp Leu His
 145 150 155 160
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 Asn Asp Leu Ile Asn Gly His Ala Val Leu Lys Glu Ile Arg Ser Ser
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 195 200 205
 Ser Thr Ala Val Gly Ala Met Ser Tyr Ala Gln Gly His Phe Ser Asn
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 Ala Phe Gly Thr Arg Ala Thr Ala Lys Ser Ala Tyr Ser Leu Ala Val
 225 230 235 240
 Gly Leu Ala Ala Thr Ala Glu Gly Gln Ser Thr Ile Ala Ile Gly Ser
 245 250 255
 Asp Ala Thr Ser Ser Ser Leu Gly Ala Ile Ala Leu Gly Ala Gly Thr
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Substitute SeqListing.txt

275

280

285

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 Ala Leu Asp Pro Lys Phe Gln Ala Thr Asn Asn Thr Lys Ala Gly Pro
 305 310 315 320
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 325 330 335
 Ala Gly Val Asn Lys Thr Asp Ala Val Asn Val Ala Gln Leu Glu Ala
 340 345 350
 Val Val Lys Trp Ala Lys Glu Arg Arg Ile Thr Phe Gln Gly Asp Asp
 355 360 365
 Asn Ser Thr Asp Val Lys Ile Gly Leu Asp Asn Thr Leu Thr Ile Lys
 370 375 380
 Gly Gly Ala Glu Thr Asn Ala Leu Thr Asp Asn Asn Ile Gly Val Val
 385 390 395 400
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 420 425 430
 Val Lys Val Gly Ser Ser Ser Ser Thr Thr Ala Glu Leu Leu Ser Asp
 435 440 445
 Ser Leu Thr Phe Thr Gln Pro Asn Thr Gly Ser Gln Ser Thr Ser Lys
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 465 470 475 480
 Thr Ala Ala Ile Gly Thr Thr Arg Ile Thr Arg Asp Lys Ile Gly Phe
 485 490 495
 Ala Arg Asp Gly Asp Val Asp Glu Lys Gln Ala Pro Tyr Leu Asp Lys
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 Lys Gln Leu Lys Val Gly Ser Val Ala Ile Thr Ile Asp Asn Gly Ile
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 Asp Ala Gly Asn Lys Lys Ile Ser Asn Leu Ala Lys Gly Ser Ser Ala
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 Ala Lys Ser Gly Asn Val Thr Ala Pro Thr Tyr Asn Ile Gly Val Lys
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Substitute SeqListing.txt

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				Ala	Leu 640
Thr 625	Val	Lys	Glu	Glu 645	Asp 650
				Ala	Asn 655
				Ala	Ile 660
Lys 625	Asp	Thr	Thr 660	Lys	Asn 665
				Ala	Gly 670
				Val	Ser 675
Gly 625	Lys	Asn 675	Gly	Leu	Thr 680
				Val	Ala 685
				Thr	Lys 690
Phe 625	Gly	Leu	Ser	Gln	Asp 695
				Asn	Gly 700
				Thr	Ile 705
Asn 625	Asn	Asp	Gly	Leu	Thr 710
				Val	Lys 715
				Asn	Glu 720
Gly 625	Ala	Asn	Gly	Ile 725	Lys 730
				Phe	Thr 735
				Thr	Asn 740
Thr 625	Gly	Ile	Ala 740	Asn	Thr 745
				Arg	Ile 750
				Thr	Arg 755
Ala 625	Gly	Ser 755	Asp	Gly	Ala 760
				Val	Asp 765
				Thr	Asn 770
Asp 625	Lys	Leu	Gln	Val	Gly 775
				Asn	Val 780
				Lys	Ile 785
Ala 625	Gly	Gly	Lys	Ala	Ile 790
				Thr	Gly 795
				Leu	Ser 800
Ala 625	Asp	Gln	Ser	Ser 805	Arg 810
				Asn	Ile 815
				Thr	Gln 820
Lys 625	Asp	Lys	Ser 820	Asn	Ala 825
				Ala	Ser 830
				Asn	Ile 835
Phe 625	Asn	Leu	Lys	Asn	Asn 840
				Pro	Ile 845
				Asp	Phe 850
Asp 625	Ile	Val	Asp	Phe	Ala 855
				Asn	Gly 860
				Ala	Thr 865
His 625	Asp	Thr	Ala	Asn	Lys 870
				Thr	Ser 875
				Lys	Val 880
Asp 625	Asp	Thr	Thr	Ile 885	His 890
				Leu	Gly 895
				Thr	Asp 900
Gly 625	Val	Lys	Thr 900	Thr	Lys 905
				Leu	Asn 910
				Ala	Ser 915
Ala 625	Thr	Asn	Phe	Asn	Val 920
				Asn	Ser 925
				Thr	Glu 930
Ala 625	Lys	Asp	Ile	Ala	Glu 935
				Leu	Asn 940
				Thr	Leu 945
Thr 625	Thr	Lys	Gly	Thr	Ala 950
				Asp	Thr 955
				Ala	Leu 960
				Gln	Thr 965
				Phe	Thr 970
				Val	Lys 975

Substitute SeqListing.txt

945 950 955 960
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 965 970 975
 Gly Gln Lys Asn Ala Asn Asn Gln Val Asn Thr Leu Thr Leu Lys Gly
 980 985 990
 Glu Asn Gly Leu Asn Ile Lys Thr Asp Lys Asn Gly Thr Val Thr Phe
 995 1000 1005
 Gly Ile Asn Thr Thr Ser Gly Leu Lys Ala Gly Lys Ser Thr Leu Asn
 1010 1015 1020
 Asp Gly Gly Leu Ser Ile Lys Asn Pro Thr Gly Ser Glu Gln Ile Gln
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 Val Gly Ala Asp Gly Val Lys Phe Ala Lys Val Asn Asn Asn Gly Val
 1045 1050 1055
 Val Gly Ala Gly Ile Asp Gly Thr Thr Arg Ile Thr Arg Asp Glu Ile
 1060 1065 1070
 Gly Phe Thr Gly Thr Asn Gly Ser Leu Asp Lys Ser Lys Pro His Leu
 1075 1080 1085
 Ser Lys Asp Gly Ile Asn Ala Gly Gly Lys Lys Ile Thr Asn Ile Gln
 1090 1095 1100
 Ser Gly Glu Ile Ala Gln Asn Ser His Asp Ala Val Thr Gly Gly Lys
 1105 1110 1115 1120
 Ile Tyr Asp Leu Lys Thr Glu Leu Glu Asn Lys Ile Ser Ser Thr Ala
 1125 1130 1135
 Lys Thr Ala Gln Asn Ser Leu His Glu Phe Ser Val Ala Asp Glu Gln
 1140 1145 1150
 Gly Asn Asn Phe Thr Val Ser Asn Pro Tyr Ser Ser Tyr Asp Thr Ser
 1155 1160 1165
 Lys Thr Ser Asp Val Ile Thr Phe Ala Gly Glu Asn Gly Ile Thr Thr
 1170 1175 1180
 Lys Val Asn Lys Gly Val Val Arg Val Gly Ile Asp Gln Thr Lys Gly
 1185 1190 1195 1200
 Leu Thr Thr Pro Lys Leu Thr Val Gly Asn Asn Asn Gly Lys Gly Ile
 1205 1210 1215
 Val Ile Asp Ser Gln Asn Gly Gln Asn Thr Ile Thr Gly Leu Ser Asn
 1220 1225 1230
 Thr Leu Ala Asn Val Thr Asn Asp Lys Gly Ser Val Arg Thr Thr Glu
 1235 1240 1245
 Gln Gly Asn Ile Ile Lys Asp Glu Asp Lys Thr Arg Ala Ala Ser Ile
 1250 1255 1260
 Val Asp Val Leu Ser Ala Gly Phe Asn Leu Gln Gly Asn Gly Glu Ala
 1265 1270 1275 1280
 Val Asp Phe Val Ser Thr Tyr Asp Thr Val Asn Phe Ala Asp Gly Asn

Substitute SeqListing.txt

1285 1290 1295
 Ala Thr Thr Ala Lys Val Thr Tyr Asp Asp Thr Ser Lys Thr Ser Lys
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 Val Val Tyr Asp Val Asn Val Asp Asp Thr Thr Ile Glu Val Lys Asp
 1315 1320 1325
 Lys Lys Leu Gly Val Lys Thr Thr Thr Leu Thr Ser Thr Gly Thr Gly
 1330 1335 1340
 Ala Asn Lys Phe Ala Leu Ser Asn Gln Ala Thr Gly Asp Ala Leu Val
 1345 1350 1355 1360
 Lys Ala Ser Asp Ile Val Ala His Leu Asn Thr Leu Ser Gly Asp Ile
 1365 1370 1375
 Gln Thr Ala Lys Gly Ala Ser Gln Ala Asn Asn Ser Ala Gly Tyr Val
 1380 1385 1390
 Asp Ala Asp Gly Asn Lys Val Ile Tyr Asp Ser Thr Asp Asn Lys Tyr
 1395 1400 1405
 Tyr Gln Ala Lys Asn Asp Gly Thr Val Asp Lys Thr Lys Glu Val Ala
 1410 1415 1420
 Lys Asp Lys Leu Val Ala Gln Ala Gln Thr Pro Asp Gly Thr Leu Ala
 1425 1430 1435 1440
 Gln Met Asn Val Lys Ser Val Ile Asn Lys Glu Gln Val Asn Asp Ala
 1445 1450 1455
 Asn Lys Lys Gln Gly Ile Asn Glu Asp Asn Ala Phe Val Lys Gly Leu
 1460 1465 1470
 Glu Lys Ala Ala Ser Asp Asn Lys Thr Lys Asn Ala Ala Val Thr Val
 1475 1480 1485
 Gly Asp Leu Asn Ala Val Ala Gln Thr Pro Leu Thr Phe Ala Gly Asp
 1490 1495 1500
 Thr Gly Thr Thr Ala Lys Lys Leu Gly Glu Thr Leu Thr Ile Lys Gly
 1505 1510 1515 1520
 Gly Gln Thr Asp Thr Asn Lys Leu Thr Asp Asn Asn Ile Gly Val Val
 1525 1530 1535
 Ala Gly Thr Asp Gly Phe Thr Val Lys Leu Ala Lys Asp Leu Thr Asn
 1540 1545 1550
 Leu Asn Ser Val Asn Ala Gly Gly Thr Lys Ile Asp Asp Lys Gly Val
 1555 1560 1565
 Ser Phe Val Asp Ser Ser Gly Gln Ala Lys Ala Asn Thr Pro Val Leu
 1570 1575 1580
 Ser Ala Asn Gly Leu Asp Leu Gly Gly Lys Val Ile Ser Asn Val Gly
 1585 1590 1595 1600
 Lys Gly Thr Lys Asp Thr Asp Ala Ala Asn Val Gln Gln Leu Asn Glu
 1605 1610 1615
 Val Arg Asn Leu Leu Gly Leu Gly Asn Ala Gly Asn Asp Asn Ala Asp

Substitute SeqListing.txt
1625 1630

1620
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1635 1640 1645
Ser Ser Ser Asn Arg Thr Val Ile Lys Ala Gly Thr Val Leu Gly Gly
1650 1655 1660
Lys Gly Asn Asn Asp Thr Glu Lys Leu Ala Thr Gly Gly Ile Gln Val
1665 1670 1675 1680
Gly Val Asp Lys Asp Gly Asn Ala Asn Gly Asp Leu Ser Asn Val Trp
1685 1690 1695
Val Lys Thr Gln Lys Asp Gly Ser Lys Lys Ala Leu Leu Ala Thr Tyr
1700 1705 1710
Asn Ala Ala Gly Gln Thr Asn Tyr Leu Thr Asn Asn Pro Ala Glu Ala
1715 1720 1725
Ile Asp Arg Ile Asn Glu Gln Gly Ile Arg Phe Phe His Val Asn Asp
1730 1735 1740
Gly Asn Gln Glu Pro Val Val Gln Gly Arg Asn Gly Ile Asp Ser Ser
1745 1750 1755 1760
Ala Ser Gly Lys His Ser Val Ala Ile Gly Phe Gln Ala Lys Ala Asp
1765 1770 1775
Gly Glu Ala Ala Val Ala Ile Gly Arg Gln Thr Gln Ala Gly Asn Gln
1780 1785 1790
Ser Ile Ala Ile Gly Asp Asn Ala Gln Ala Thr Gly Asp Gln Ser Ile
1795 1800 1805
Ala Ile Gly Thr Gly Asn Val Val Ala Gly Lys His Ser Gly Ala Ile
1810 1815 1820
Gly Asp Pro Ser Thr Val Lys Ala Asp Asn Ser Tyr Ser Val Gly Asn
1825 1830 1835 1840
Asn Asn Gln Phe Thr Asp Ala Thr Gln Thr Asp Val Phe Gly Val Gly
1845 1850 1855
Asn Asn Ile Thr Val Thr Glu Ser Asn Ser Val Ala Leu Gly Ser Asn
1860 1865 1870
Ser Ala Ile Ser Ala Gly Thr His Ala Gly Thr Gln Ala Lys Lys Ser
1875 1880 1885
Asp Gly Thr Ala Gly Thr Thr Thr Thr Ala Gly Ala Thr Gly Thr Val
1890 1895 1900
Lys Gly Phe Ala Gly Gln Thr Ala Val Gly Ala Val Ser Val Gly Ala
1905 1910 1915 1920
Ser Gly Ala Glu Arg Arg Ile Gln Asn Val Ala Ala Gly Glu Val Ser
1925 1930 1935
Ala Thr Ser Thr Asp Ala Val Asn Gly Ser Gln Leu Tyr Lys Ala Thr
1940 1945 1950
Gln Ser Ile Ala Asn Ala Thr Asn Glu Leu Asp His Arg Ile His Gln

Substitute SeqListing.txt

1955

1960

1965

Asn Glu Asn Lys Ala Asn Ala Gly Ile Ser Ser Ala Met Ala Met Ala
1970 1975 1980

Ser Met Pro Gln Ala Tyr Ile Pro Gly Arg Ser Met Val Thr Gly Gly
1985 1990 1995 2000

Ile Ala Thr His Asn Gly Gln Gly Ala Val Ala Val Gly Leu Ser Lys
2005 2010 2015

Leu Ser Asp Asn Gly Gln Trp Val Phe Lys Ile Asn Gly Ser Ala Asp
2020 2025 2030

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2035 2040 2045

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<212> DNA

<213> Moraxella catarrhalis

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Substitute SeqListing.txt

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130 135 140Val Leu Ala Glu Gly Asp Ala Ser Ile Ala Ile Gly Ser Asp Asp Leu
145 150 155 160Tyr Leu Pro Lys Asn Leu Asp Leu Lys Asn Glu Phe His Lys Leu Ile
165 170 175His Gly His Glu Ile Leu Lys Lys Ile Gln Thr Ser Thr Asp Gly Lys
180 185 190Ile Lys Tyr Arg Arg Thr Arg Ala Gln Gly His Ala Ser Thr Ala Val
195 200 205Gly Ala Met Ser Tyr Ala Gln Gly His Phe Ser Asn Ala Phe Gly Thr
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245 250 255Asn Ala Phe Ala Ala Thr Ala Ile Gly Gly Asn Thr Val Val Asn Leu
260 265 270Gly Arg Gly Val Ala Leu Gly Phe Gly Ser Gln Ile Leu Asp Arg Asp
275 280 285

Substitute SeqListing.txt

Asn 290 Thr Asp Ala Ser 295 Tyr Val Pro Leu 300 Gly Lys Thr Leu Ala
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 Asn Val Gly 340 Ala Gly Ser Arg Asp Thr 345 Asp Ala Val Asn 350 Val Ala Gln
 Leu Lys 355 Leu Val Glu Glu Leu 360 Ala Asn Arg Lys Ile Thr 365 Phe Lys Gly
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 Ile Gly Val Val Thr 405 Asp Gly Asn Gly Leu 410 Lys Val Lys Leu Ala 415 Lys
 Glu Leu Thr 420 Gly Leu Thr Ser Val 425 Ala Thr Asn Lys 430 Ile Thr Val
 Ser Asn Thr 435 Asn Asn Asn Asn Ala 440 Glu Leu Gln Ser Gly 445 Gly Leu Thr
 Phe Ser 450 Pro Ile Thr Gly Thr 455 Lys Thr Asp Lys 460 Thr Val Tyr Ser Ile
 Asp 465 Gly Leu Lys Phe Thr 470 Asn Asp Ser Asn Ser 475 Ile Ala Thr Lys Gly 480
 Thr Thr Arg Ile Thr 485 Lys Lys Lys Ile Gly 490 Phe Ala Gly Thr Asn 495 Asp
 Gly Val Asp 500 Glu Ser Lys Pro Tyr Leu 505 Asp Asn Glu Lys Leu 510 Lys Val
 Gly Asn Ser 515 Thr Leu Asn Ser Gly 520 Ser Leu Thr Val Asn 525 Asn Thr Thr
 Gly Asn 530 Lys Gln Ile Gln Val 535 Gly Ala Asn Gly Ile 540 Lys Phe Ala Thr
 Val 545 Ala Asn Asn Val Ala 550 Asn Thr Ser Ala Thr 555 Val Gly Thr Ala Arg 560
 Ile Thr Glu Glu Lys 565 Ile Gly Phe Ala Gly 570 Thr Asn Asp Gly Val 575 Asp
 Glu Gln Ala Pro 580 Tyr Leu Asp Lys Glu 585 Arg Leu Lys Val Gly 590 Arg Val
 Glu Ile Thr 595 Thr Asp Ser Gly Ile 600 Asn Ala Gly Asn His 605 Lys Ile Thr
 Gly Leu 610 Thr Asn Gly Ile Ala 615 Asn Thr Asp Ala Val 620 Thr Ile Lys Gln

Substitute SeqListing.txt

Leu Lys Asp Ala Lys Pro Thr Leu Asn Ala Gly Asp Gly Ile Ser Ile
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 Asn Ser Asn Asn Gly Asp Leu Val Asp Ser Ser Gly Asn Ile Thr Thr
 645 650 655
 Pro Thr Tyr Asn Ile Ser Val Lys Thr Thr Lys Leu Asn Ser Asn Gly
 660 665 670
 Thr Ser Gly Asn Asn Lys Phe Ser Val Ser Asn Ala His Asp Asn Asn
 675 680 685
 Ser Leu Val Thr Ala Lys Asp Leu Ala Asp Tyr Leu Asn Lys Val Asn
 690 695 700
 Glu Thr Ala Asp Ser Ala Leu Pro Ser Phe Lys Val Gln Asn Gly Asp
 705 710 715 720
 Asn Ser Asn Asn Ala Ile Thr Val Gly Lys Asp Thr Asn Gly Lys Thr
 725 730 735
 Phe Asn Thr Leu Lys Leu Lys Gly Glu Asn Gly Val Asn Ile Thr Thr
 740 745 750
 Asn Arg Ala Thr Gly Thr Val Thr Phe Gly Ile Asp Gln Ser Asn Gly
 755 760 765
 Leu Thr Thr Pro Lys Leu Thr Val Gly Ser Asp Thr Asn Gly Asn Arg
 770 775 780
 Leu Val Ile Glu Gln Val Pro Ser Ala Asp Gly Asn Ser Thr Lys Asn
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 Ile Ile Lys Gly Leu Ser Pro Thr Leu Pro Ser Ile Ala Ser Pro Ser
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 Gly Arg Asn Ile Ala Leu Gly Asn Thr Ile Glu Glu Lys Asp Lys Ser
 820 825 830
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 850 855 860
 Phe Ile Asp Gly Asn Ala Thr Thr Ala Thr Val Thr Tyr Asp Glu Ala
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 Asn Gln Thr Ser Lys Val Ala Tyr Asp Val Asn Val Asp Glu Lys Thr
 885 890 895
 Ile Glu Leu Thr Gly Asp Asn Gly Lys Lys Gln Leu Gly Val Lys Thr
 900 905 910
 Ile Lys Leu Thr Glu Thr Ser Thr Asn Gly Asn Ala Thr Thr Phe Ser
 915 920 925
 Thr Asp Asp Asp His Ala Leu Val Lys Ala Ser Asp Ile Ala Gly Asn
 930 935 940
 Leu Asn Thr Leu Ala Glu Glu Ile His Thr Thr Lys Gly Thr Ala Asn
 945 950 955 960

Substitute SeqListing.txt

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Ala Asp Asp Thr Asn Ala Ile Thr Val Gly Lys Asp Gly Thr Ser Gly
980 985 990

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995 1000 1005

Thr Asp Lys Asp Gly Thr Val Thr Phe Gly Ile Asn Thr Gln Ser Gly
1010 1015 1020

Leu Lys Ala Gly Asp Ser Thr Thr Leu Asn Asn Asn Gly Leu Ser Ile
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Lys Asn Thr Ala Ser Asn Glu Gln Ile Gln Val Gly Ala Asp Gly Val
1045 1050 1055

Lys Phe Ala Met Val Asn Asn Gly Val Val Gly Ala Gly Ile Asp Gly
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Thr Thr Arg Ile Thr Arg Asp Glu Ile Gly Phe Thr Gly Thr Asn Gly
1075 1080 1085

Ser Leu Asp Lys Ser Lys Pro His Leu Ser Lys Asp Gly Ile Asn Ala
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Gly Gly Lys Lys Ile Thr Asn Ile Gln Ser Gly Glu Ile Ala Lys Asn
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Ser His Asp Ala Val Thr Gly Gly Lys Ile Tyr Asp Leu Lys Thr Glu
1125 1130 1135

Leu Glu Asn Lys Ile Ser Ser Thr Ala Lys Thr Ala Gln Asn Ser Leu
1140 1145 1150

His Glu Phe Ser Val Ala Asp Glu Gln Gly Asn Asn Phe Thr Val Ser
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Asn Pro Tyr Ser Ser Tyr Asp Thr Ser Lys Thr Ser Asp Val Ile Thr
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Phe Ala Gly Glu Asn Gly Ile Thr Thr Lys Val Asn Lys Gly Val Val
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Arg Val Gly Ile Asp Gln Thr Lys Gly Leu Thr Thr Pro Lys Leu Thr
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Gln Asn Thr Ile Thr Gly Leu Ser Asn Thr Leu Ala Asn Val Thr Asn
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Asp Lys Gly Ser Val Arg Thr Thr Glu Gln Gly Asn Ile Ile Lys Asp
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Glu Asp Lys Thr Arg Ala Ser Ile Val Asp Val Leu Ser Ala Gly
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Phe Asn Leu Gln Gly Asn Gly Glu Ala Val Asp Phe Val Ser Thr Tyr
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Substitute SeqListing.txt

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Tyr Asp Asp Thr Ser Lys Thr Ser Lys Val Val Tyr Asp Val Asn Val
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Asp Asp Thr Thr Ile Glu Val Lys Asp Lys Lys Leu Gly Val Lys Thr
1330 1335 1340

Thr Thr Leu Thr Ser Thr Gly Thr Gly Ala Asn Lys Phe Ala Leu Ser
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Asn Gln Ala Thr Gly Asp Ala Leu Val Lys Ala Ser Asp Ile Val Ala
1365 1370 1375

His Leu Asn Thr Leu Ser Gly Asp Ile Gln Thr Ala Lys Gly Ala Ser
1380 1385 1390

Gln Ala Asn Asn Ser Ala Gly Tyr Val Asp Ala Asp Gly Asn Lys Val
1395 1400 1405

Ile Tyr Asp Ser Thr Asp Asn Lys Tyr Tyr Gln Ala Lys Asn Asp Gly
1410 1415 1420

Thr Val Asp Lys Thr Lys Glu Val Ala Lys Asp Lys Leu Val Ala Gln
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Ala Gln Thr Pro Asp Gly Thr Leu Ala Gln Met Asn Val Lys Ser Val
1445 1450 1455

Ile Asn Lys Glu Gln Val Asn Asp Ala Asn Lys Lys Gln Gly Ile Asn
1460 1465 1470

Glu Asp Asn Ala Phe Val Lys Gly Leu Glu Lys Ala Ala Ser Asp Asn
1475 1480 1485

Lys Thr Lys Asn Ala Ala Val Thr Val Gly Asp Leu Asn Ala Val Ala
1490 1495 1500

Gln Thr Pro Leu Thr Phe Ala Gly Asp Thr Gly Thr Thr Ala Lys Lys
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Leu Gly Glu Thr Leu Thr Ile Lys Gly Gly Gln Thr Asp Thr Asn Lys
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Leu Thr Asp Asn Asn Ile Gly Val Val Ala Gly Thr Asp Gly Phe Thr
1540 1545 1550

Val Lys Leu Ala Lys Asp Leu Thr Asn Leu Asn Ser Val Asn Ala Gly
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Gly Thr Lys Ile Asp Glu Lys Gly Ile Ser Phe Val Asp Ala Asn Gly
1570 1575 1580

Gln Ala Lys Ala Asn Thr Pro Val Leu Ser Ala Asn Gly Leu Asp Leu
1585 1590 1595 1600

Gly Gly Lys Val Ile Ser Asn Val Gly Lys Gly Thr Lys Asp Thr Asp
1605 1610 1615

Ala Ala Asn Val Gln Gln Leu Asn Glu Val Arg Asn Leu Leu Gly Leu
1620 1625 1630

Substitute SeqListing.txt

Gly Asn Asp Asn Ala Asp Gly Asn Gln Val Asn Ile Ala Asp Ile Lys
1635 1640 1645

Lys Asp Pro Asn Ser Gly Ser Ser Ser Asn Arg Thr Val Ile Lys Ala
1650 1655 1660

Gly Thr Val Leu Gly Gly Lys Gly Asn Asn Asp Thr Glu Lys Leu Ala
1665 1670 1675 1680

Thr Gly Gly Val Gln Val Gly Val Asp Lys Asp Gly Asn Ala Asn Gly
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Asp Leu Ser Asn Val Trp Val Lys Thr Gln Lys Asp Gly Ser Lys Lys
1700 1705 1710

Ala Leu Leu Ala Thr Tyr Asn Ala Ala Gly Gln Thr Asn Tyr Val Thr
1715 1720 1725

Asn Asn Pro Ala Glu Ala Ile Asp Arg Ile Asn Glu Gln Gly Ile Arg
1730 1735 1740

Phe Phe His Val Asn Asp Gly Asn Gln Glu Pro Val Val Gln Gly Arg
1745 1750 1755 1760

Asn Gly Ile Asp Ser Ser Ala Ser Gly Lys His Ser Val Ala Ile Gly
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Phe Gln Ala Lys Ala Asp Gly Glu Ala Ala Val Ala Ile Gly Arg Gln
1780 1785 1790

Thr Gln Ala Gly Asn Gln Ser Ile Ala Ile Gly Asp Asn Ala Gln Ala
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1810 1815 1820

Lys His Ser Gly Ala Ile Gly Asp Pro Ser Thr Val Lys Ala Asp Asn
1825 1830 1835 1840

Ser Tyr Ser Val Gly Asn Asn Asn Gln Phe Thr Asp Ala Thr Gln Thr
1845 1850 1855

Asp Val Phe Gly Val Gly Asn Asn Ile Thr Val Thr Glu Ser Asn Ser
1860 1865 1870

Val Ala Leu Gly Ser Asn Ser Ala Ile Ser Ala Gly Thr His Ala Gly
1875 1880 1885

Thr Gln Ala Lys Lys Ser Asp Gly Thr Ala Gly Thr Thr Thr Ala
1890 1895 1900

Gly Ala Thr Gly Thr Val Lys Gly Phe Ala Gly Gln Thr Ala Val Gly
1905 1910 1915 1920

Ala Val Ser Val Gly Ala Ser Gly Ala Glu Arg Arg Ile Gln Asn Val
1925 1930 1935

Ala Ala Gly Glu Val Ser Ala Thr Ser Thr Asp Ala Val Asn Gly Ser
1940 1945 1950

Gln Leu Tyr Lys Ala Thr Gln Ser Ile Ala Asn Ala Thr Asn Glu Leu
1955 1960 1965

Substitute SeqListing.txt

Asp His Arg Ile His Gln Asn Glu Asn Lys Ala Asn Ala Gly Ile Ser
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Ser Ala Met Ala Met Ala Ser Met Pro Gln Ala Tyr Ile Pro Gly Arg
1985 1990 1995 2000

Ser Met Val Thr Gly Gly Ile Ala Thr His Asn Gly Gln Gly Ala Val
2005 2010 2015

Ala Val Gly Leu Ser Lys Leu Ser Asp Asn Gly Gln Trp Val Phe Lys
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Substitute SeqListing.txt

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Substitute SeqListing.txt

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Substitute SeqListing.txt

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Substitute SeqListing.txt

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Thr Asp 610 Asp Thr Asp Ala Val 615 Thr Tyr Lys Gln Leu 620 Lys Gln Val Gln

Gln Asp Ala Asp Gly 630 Ala Leu Gln Ser Phe 635 Ser Ile Arg Asp Glu Lys 640

Gly Gln Glu Phe Thr 645 Ile Ser Asn Leu Tyr 650 Ser Asn Gly Asn Thr Pro

Asn Thr Phe 660 Glu Thr Ile Thr Phe 665 Ala Gly Glu Asn Gly Ile 670 Ser Ile

Ser Asn Asp 675 Ile Ala Lys Gly Lys 680 Val Lys Val Gly Ile 685 Asp Pro Ile

Asn Gly 690 Leu Thr Thr Pro Lys 695 Leu Thr Val Gly Ser 700 Asp Lys Asp Gly

Lys Thr Gln Leu Val Ile 710 Glu Gln Val Ala Ser 715 Gly Asn Asp Thr Lys 720

Asn Ile Ile Arg Gly 725 Leu Ser Pro Thr Leu 730 Pro Ser Ile Thr Asn Ala 735

Gly Gly Val Arg 740 Thr Thr Glu Gln Gly 745 Asn Thr Ile Thr Ser Asp Glu 750

Asp Lys Ser 755 Lys Ala Ala Ser Ile 760 Gly Asp Ile Leu Asn 765 Thr Gly Phe

Asn Leu 770 Lys Asn Asn Ser Asn 775 Ser Val Gly Phe Val 780 Ser Thr Tyr Asn

Thr Val Asp Phe Ile Asp 790 Gly Asn Ala Thr Thr 795 Ala Lys Val Thr Tyr 800

Asp Glu Thr Asn 805 Gln Thr Ser Lys Val Thr 810 Tyr Asp Val Asn Val 815 Asp

Glu Lys Thr Ile 820 Glu Leu Thr Gly Asp 825 Asn Gly Lys Thr Asn Lys Ile 830

Gly Val Lys 835 Thr Thr Thr Leu Thr 840 Thr Thr Asn Ala Asn 845 Gly Lys Ala

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Substitute SeqListing.txt

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895

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965 970 975
Gly Val Lys Phe Ala Lys Val Asp Lys Gly Asn Ser Ser Thr Gly Ile
980 985 990
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995 1000 1005
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1125 1130 1135
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Substitute SeqListing.txt

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 Asp Gly Thr Leu Ala Gln Met Asn Val Lys Ser Val Ile Asn Lys Glu
 1380 1385 1390
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 Thr Phe Ala Gly Asp Thr Gly Thr Thr Ala Lys Lys Leu Gly Glu Thr
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 Leu Thr Ile Lys Gly Gly Gln Thr Asp Thr Asn Lys Leu Thr Asp Asn
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 Asn Ile Gly Val Val Ala Gly Thr Asp Gly Phe Thr Val Lys Leu Ala
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 Lys Asp Leu Thr Asn Leu Asn Ser Val Asn Ala Gly Gly Thr Arg Ile
 1490 1495 1500
 Asp Glu Lys Gly Ile Ser Phe Val Asp Ala Asn Gly Gln Ala Lys Ala
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 Ile Ser Asn Ile Gly Ala Ala Val Asp Asp Asn Asp Ala Val Asn Phe
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Substitute SeqListing.txt

1555

1560

1565

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Ala His Gly Lys Pro Leu Asp Ala Gly His Gln Val Val Ala Ser Leu
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Gly Gly Asn Ser Asp Ala Ile Thr Leu Thr Asn Ile Lys Ser Thr Leu
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Pro Gln Ile Asp Thr Pro Asn Thr Gly Asn Ala Asn Ala Gly Gln Ala
1665 1670 1675 1680

Gln Ser Leu Pro Ser Leu Ser Ala Ala Gln Gln Ser Asn Ala Ala Ser
1685 1690 1695

Val Lys Asp Val Leu Asn Val Gly Phe Asn Leu Gln Thr Asn His Asn
1700 1705 1710

Gln Val Asp Phe Val Lys Ala Tyr Asp Thr Val Asn Phe Val Asn Gly
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1745 1750 1755 1760

Val Leu Ile Lys Ala Lys Asp Gly Lys Phe Tyr Lys Ala Asp Asp Leu
1765 1770 1775

Met Pro Asn Gly Ser Leu Lys Ala Gly Lys Ser Ala Ser Asp Ala Lys
1780 1785 1790

Thr Pro Thr Gly Leu Ser Leu Val Asn Pro Asn Ala Gly Lys Gly Ser
1795 1800 1805

Thr Gly Asp Ala Val Ala Leu Asn Asn Leu Ser Lys Ala Val Phe Lys
1810 1815 1820

Ser Lys Asp Gly Thr Thr Thr Thr Thr Val Ser Ser Asp Gly Ile Ser
1825 1830 1835 1840

Ile Gln Gly Lys Asp Asn Ser Ser Ile Thr Leu Ser Lys Asp Gly Leu
1845 1850 1855

Asn Val Gly Gly Lys Val Ile Ser Asn Val Gly Lys Gly Thr Lys Asp
1860 1865 1870

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1875 1880 1885

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Substitute SeqListing.txt

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1955 1960 1965
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Substitute SeqListing.txt

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Substitute SeqListing.txt

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 Val Asn Gly Ser Ser Leu Asp Lys Ile Gly Thr Asp Ala Thr Gly Gln
 65 70 75 80
 Glu Ser Ile Ala Ile Gly Gly Asp Val Lys Ala Ser Gly Asp Ala Ser
 85 90 95
 Ile Ala Ile Gly Ser Asp Asp Leu His Leu Leu Asp Gln His Gly Asn
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 Ser Tyr Ala Gln Gly His Phe Ser Asn Ala Phe Gly Thr Arg Ala Thr
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 225 230 235 240
 Arg Pro Ala Tyr Thr Pro Asn Thr Gln Ala Leu Asp Pro Lys Phe Gln
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 260 265 270
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Substitute SeqListing.txt
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325 330 335
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340 345 350
Leu Lys Val Lys Leu Ala Lys Thr Leu Asn Asn Leu Thr Glu Val Asn
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450 455 460
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465 470 475 480
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485 490 495
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Ala Pro Thr Tyr Asn Ile Gly Val Lys Thr Thr Glu Leu Asn Ser Asp
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Thr Ala Asp Ser Ala Leu Gln Ser Phe Thr Val Lys Glu Glu Asp Asp
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Substitute SeqListing.txt

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725 730 735

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755 760 765

Ser Ile Asn Asp Ile Leu Asn Thr Gly Phe Asn Leu Lys Asn Asn Asn
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Thr Gly Thr Asp Asp Asn Lys Lys Leu Gly Val Lys Thr Thr Lys Leu
835 840 845

Asn Lys Thr Ser Ala Asn Gly Asn Thr Ala Thr Asn Phe Asn Val Asn
850 855 860

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Substitute SeqListing.txt

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 995 1000 1005
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 Ser His Asp Ala Val Thr Gly Gly Lys Ile Tyr Asp Leu Lys Thr Glu
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 Leu Glu Asn Lys Ile Ser Ser Thr Ala Lys Thr Ala Gln Asn Ser Leu
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 His Glu Phe Ser Val Ala Asp Glu Gln Gly Asn Asn Phe Thr Val Ser
 1090 1095 1100
 Asn Pro Tyr Ser Ser Tyr Asp Thr Ser Lys Thr Ser Asp Val Ile Thr
 1105 1110 1115 1120
 Phe Ala Gly Glu Asn Gly Ile Thr Thr Lys Val Asn Lys Gly Val Val
 1125 1130 1135
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 Tyr Asp Asp Thr Ser Lys Thr Ser Lys Val Val Tyr Asp Val Asn Val
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Substitute SeqListing.txt

1285

1290

1295

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1315 1320 1325

Gln Ala Asn Asn Ser Ala Gly Tyr Val Asp Ala Asp Gly Asn Lys Val
1330 1335 1340

Ile Tyr Asp Ser Thr Asp Asn Lys Tyr Tyr Gln Ala Lys Asn Asp Gly
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Thr Val Asp Lys Thr Lys Glu Val Ala Lys Asp Lys Leu Val Ala Gln
1365 1370 1375

Ala Gln Thr Pro Asp Gly Thr Leu Ala Gln Met Asn Val Lys Ser Val
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Ile Asn Lys Glu Gln Val Asn Asp Ala Asn Lys Lys Gln Gly Ile Asn
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Lys Thr Lys Asn Ala Ala Val Thr Val Gly Asp Leu Asn Ala Val Ala
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Leu Gly Glu Thr Leu Thr Ile Lys Gly Gly Gln Thr Asp Thr Asn Lys
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Val Lys Leu Ala Lys Asp Leu Thr Asn Leu Asn Ser Val Asn Ala Gly
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Gly Thr Lys Ile Asp Asp Lys Gly Val Ser Phe Val Asp Ser Ser Gly
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Ile Lys Ala Gly Thr Val Leu Gly Gly Lys Gly Asn Asn Asp Thr Glu
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Lys Leu Ala Thr Gly Gly Ile Gln Val Gly Val Asp Lys Asp Gly Asn
Page 50

Substitute SeqListing.txt

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 1925 1930 1935
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 1940 1945 1950
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Substitute SeqListing.txt

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Substitute SeqListing.txt

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Substitute SeqListing.txt

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Substitute SeqListing.txt

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Substitute SeqListing.txt

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```

<210> 46

<211> 1044

<212> PRT

<213> Moraxella catarrhalis

<400> 46

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Lys Ser Thr Leu Asn Asp Gly Gly Leu Ser Ile Lys Asn Pro Thr Gly
20 25 30

Ser Glu Gln Ile Gln Val Gly Ala Asp Gly Val Lys Phe Ala Lys Val
35 40 45

Asn Asn Asn Gly Val Val Gly Ala Gly Ile Asp Gly Thr Thr Arg Ile
50 55 60

Thr Arg Asp Glu Ile Gly Phe Thr Gly Thr Asn Gly Ser Leu Asp Lys
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Substitute SeqListing.txt

65 70 75 80

Ser Lys Pro His Leu Ser Lys Asp Gly Ile Asn Ala Gly Gly Lys Lys
 85 90 95

Ile Thr Asn Ile Gln Ser Gly Glu Ile Ala Gln Asn Ser His Asp Ala
 100 105 110

Val Thr Gly Gly Lys Ile Tyr Asp Leu Lys Thr Glu Leu Glu Asn Lys
 115 120 125

Ile Ser Ser Thr Ala Lys Thr Ala Gln Asn Ser Leu His Glu Phe Ser
 130 135 140

Val Ala Asp Glu Gln Gly Asn Asn Phe Thr Val Ser Asn Pro Tyr Ser
 145 150 155 160

Ser Tyr Asp Thr Ser Lys Thr Ser Asp Val Ile Thr Phe Ala Gly Glu
 165 170 175

Asn Gly Ile Thr Thr Lys Val Asn Lys Gly Val Val Arg Val Gly Ile
 180 185 190

Asp Gln Thr Lys Gly Leu Thr Thr Pro Lys Leu Thr Val Gly Asn Asn
 195 200 205

Asn Gly Lys Gly Ile Val Ile Asp Ser Gln Asn Gly Gln Asn Thr Ile
 210 215 220

Thr Gly Leu Ser Asn Thr Leu Ala Asn Val Thr Asn Asp Lys Gly Ser
 225 230 235 240

Val Arg Thr Thr Glu Gln Gly Asn Ile Ile Lys Asp Glu Asp Lys Thr
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Arg Ala Ala Ser Ile Val Asp Val Leu Ser Ala Gly Phe Asn Leu Gln
 260 265 270

Gly Asn Gly Glu Ala Val Asp Phe Val Ser Thr Tyr Asp Thr Val Asn
 275 280 285

Phe Ala Asp Gly Asn Ala Thr Thr Ala Lys Val Thr Tyr Asp Asp Thr
 290 295 300

Ser Lys Thr Ser Lys Val Val Tyr Asp Val Asn Val Asp Asp Thr Thr
 305 310 315 320

Ile Glu Val Lys Asp Lys Lys Leu Gly Val Lys Thr Thr Thr Leu Thr
 325 330 335

Ser Thr Gly Thr Gly Ala Asn Lys Phe Ala Leu Ser Asn Gln Ala Thr
 340 345 350

Gly Asp Ala Leu Val Lys Ala Ser Asp Ile Val Ala His Leu Asn Thr
 355 360 365

Leu Ser Gly Asp Ile Gln Thr Ala Lys Gly Ala Ser Gln Ala Asn Asn
 370 375 380

Ser Ala Gly Tyr Val Asp Ala Asp Gly Asn Lys Val Ile Tyr Asp Ser
 385 390 395 400

Thr Asp Asn Lys Tyr Tyr Gln Ala Lys Asn Asp Gly Thr Val Asp Lys

Substitute SeqListing.txt

405 410 415
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 420 425 430
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 435 440 445
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 450 455 460
 Phe Val Lys Gly Leu Glu Lys Ala Ala Ser Asp Asn Lys Thr Lys Asn
 465 470 475 480
 Ala Ala Val Thr Val Gly Asp Leu Asn Ala Val Ala Gln Thr Pro Leu
 485 490 495
 Thr Phe Ala Gly Asp Thr Gly Thr Thr Ala Lys Lys Leu Gly Glu Thr
 500 505 510
 Leu Thr Ile Lys Gly Gly Gln Thr Asp Thr Asn Lys Leu Thr Asp Asn
 515 520 525
 Asn Ile Gly Val Val Ala Gly Thr Asp Gly Phe Thr Val Lys Leu Ala
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 Asp Asp Lys Gly Val Ser Phe Val Asp Ser Ser Gly Gln Ala Lys Ala
 565 570 575
 Asn Thr Pro Val Leu Ser Ala Asn Gly Leu Asp Leu Gly Gly Lys Val
 580 585 590
 Ile Ser Asn Val Gly Lys Gly Thr Lys Asp Thr Asp Ala Ala Asn Val
 595 600 605
 Gln Gln Leu Asn Glu Val Arg Asn Leu Leu Gly Leu Gly Asn Ala Gly
 610 615 620
 Asn Asp Asn Ala Asp Gly Asn Gln Val Asn Ile Ala Asp Ile Lys Lys
 625 630 635 640
 Asp Pro Asn Ser Gly Ser Ser Ser Asn Arg Thr Val Ile Lys Ala Gly
 645 650 655
 Thr Val Leu Gly Gly Lys Gly Asn Asn Asp Thr Glu Lys Leu Ala Thr
 660 665 670
 Gly Gly Ile Gln Val Gly Val Asp Lys Asp Gly Asn Ala Asn Gly Asp
 675 680 685
 Leu Ser Asn Val Trp Val Lys Thr Gln Lys Asp Gly Ser Lys Lys Ala
 690 695 700
 Leu Leu Ala Thr Tyr Asn Ala Ala Gly Gln Thr Asn Tyr Leu Thr Asn
 705 710 715 720
 Asn Pro Ala Glu Ala Ile Asp Arg Ile Asn Glu Gln Gly Ile Arg Phe
 725 730 735
 Phe His Val Asn Asp Gly Asn Gln Glu Pro Val Val Gln Gly Arg Asn

Substitute SeqListing.txt

740
 Gly Ile Asp Ser Ser Ala Ser Gly Lys His Ser Val Ala Ile Gly Phe
 755 760 765
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 770 775 780
 Gln Ala Gly Asn Gln Ser Ile Ala Ile Gly Asp Asn Ala Gln Ala Thr
 785 790 795
 Gly Asp Gln Ser Ile Ala Ile Gly Thr Gly Asn Val Val Ala Gly Lys
 805 810 815
 His Ser Gly Ala Ile Gly Asp Pro Ser Thr Val Lys Ala Asp Asn Ser
 820 825 830
 Tyr Ser Val Gly Asn Asn Asn Gln Phe Thr Asp Ala Thr Gln Thr Asp
 835 840 845
 Val Phe Gly Val Gly Asn Asn Ile Thr Val Thr Glu Ser Asn Ser Val
 850 855 860
 Ala Leu Gly Ser Asn Ser Ala Ile Ser Ala Gly Thr His Ala Gly Thr
 865 870 875 880
 Gln Ala Lys Lys Ser Asp Gly Thr Ala Gly Thr Thr Thr Thr Ala Gly
 885 890 895
 Ala Thr Gly Thr Val Lys Gly Phe Ala Gly Gln Thr Ala Val Gly Ala
 900 905 910
 Val Ser Val Gly Ala Ser Gly Ala Glu Arg Arg Ile Gln Asn Val Ala
 915 920 925
 Ala Gly Glu Val Ser Ala Thr Ser Thr Asp Ala Val Asn Gly Ser Gln
 930 935 940
 Leu Tyr Lys Ala Thr Gln Ser Ile Ala Asn Ala Thr Asn Glu Leu Asp
 945 950 955 960
 His Arg Ile His Gln Asn Glu Asn Lys Ala Asn Ala Gly Ile Ser Ser
 965 970 975
 Ala Met Ala Met Ala Ser Met Pro Gln Ala Tyr Ile Pro Gly Arg Ser
 980 985 990
 Met Val Thr Gly Gly Ile Ala Thr His Asn Gly Gln Gly Ala Val Ala
 995 1000 1005
 Val Gly Leu Ser Lys Leu Ser Asp Asn Gly Gln Trp Val Phe Lys Ile
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 Asn Gly Ser Ala Asp Thr Gln Gly His Val Gly Ala Ala Val Gly Ala
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 Gly Phe His Phe

<210> 47
 <211> 2448
 <212> DNA

Substitute SeqListing.txt

<213> Moraxella catarrhalis

<400> 47

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<210> 48

<211> 816

<212> PRT

<213> Moraxella catarrhalis

<400> 48

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Ser Glu Gln Ile Gln Val Gly Ala Asp Gly Val Lys Phe Ala Lys Val
          35          40          45
Asn Asn Asn Gly Val Val Gly Ala Gly Ile Asp Gly Thr Thr Arg Ile
          50          55          60

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Substitute SeqListing.txt

Thr Arg Asp Glu Ile Gly Phe Thr Gly Thr Asn Gly Ser Leu Asp Lys
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 85 90 95
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 100 105 110
 Val Thr Gly Gly Lys Ile Tyr Asp Leu Lys Thr Glu Leu Glu Asn Lys
 115 120 125
 Ile Ser Ser Thr Ala Lys Thr Ala Gln Asn Ser Leu His Glu Phe Ser
 130 135 140
 Val Ala Asp Glu Gln Gly Asn Asn Phe Thr Val Ser Asn Pro Tyr Ser
 145 150 155 160
 Ser Tyr Asp Thr Ser Lys Thr Ser Asp Val Ile Thr Phe Ala Gly Glu
 165 170 175
 Asn Gly Ile Thr Thr Lys Val Asn Lys Gly Val Val Arg Val Gly Ile
 180 185 190
 Asp Gln Thr Lys Gly Leu Thr Thr Pro Lys Leu Thr Val Gly Asn Asn
 195 200 205
 Asn Gly Lys Gly Ile Val Ile Asp Ser Gln Asn Gly Gln Asn Thr Ile
 210 215 220
 Thr Gly Leu Ser Asn Thr Leu Ala Asn Val Thr Asn Asp Lys Gly Ser
 225 230 235 240
 Val Arg Thr Thr Glu Gln Gly Asn Ile Ile Lys Asp Glu Asp Lys Thr
 245 250 255
 Arg Ala Ala Ser Ile Val Asp Val Leu Ser Ala Gly Phe Asn Leu Gln
 260 265 270
 Gly Asn Gly Glu Ala Val Asp Phe Val Ser Thr Tyr Asp Thr Val Asn
 275 280 285
 Phe Ala Asp Gly Asn Ala Thr Thr Ala Lys Val Thr Tyr Asp Asp Thr
 290 295 300
 Ser Lys Thr Ser Lys Val Val Tyr Asp Val Asn Val Asp Asp Thr Thr
 305 310 315 320
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 325 330 335
 Ser Thr Gly Thr Gly Ala Asn Lys Phe Ala Leu Ser Asn Gln Ala Thr
 340 345 350
 Gly Asp Ala Leu Val Lys Ala Ser Asp Ile Val Ala His Leu Asn Thr
 355 360 365
 Leu Ser Gly Asp Ile Gln Thr Ala Lys Gly Ala Ser Gln Ala Asn Asn
 370 375 380
 Ser Ala Gly Tyr Val Asp Ala Asp Gly Asn Lys Val Ile Tyr Asp Ser
 385 390 395 400

Substitute SeqListing.txt

Thr Asp Asn Lys Tyr Tyr Gln Ala Lys Asn Asp Gly Thr Val Asp Lys
405 410 415

Thr Lys Glu Val Ala Lys Asp Lys Leu Val Ala Gln Ala Gln Thr Pro
420 425 430

Asp Gly Thr Leu Ala Gln Met Asn Val Lys Ser Val Ile Asn Lys Glu
435 440 445

Gln Val Asn Asp Ala Asn Lys Lys Gln Gly Ile Asn Glu Asp Asn Ala
450 455 460

Phe Val Lys Gly Leu Glu Lys Ala Ala Ser Asp Asn Lys Thr Lys Asn
465 470 475 480

Ala Ala Val Thr Val Gly Asp Leu Asn Ala Val Ala Gln Thr Pro Leu
485 490 495

Thr Phe Ala Gly Asp Thr Gly Thr Thr Ala Lys Lys Leu Gly Glu Thr
500 505 510

Leu Thr Ile Lys Gly Gly Gln Thr Asp Thr Asn Lys Leu Thr Asp Asn
515 520 525

Asn Ile Gly Val Val Ala Gly Thr Asp Gly Phe Thr Val Lys Leu Ala
530 535 540

Lys Asp Leu Thr Asn Leu Asn Ser Val Asn Ala Gly Gly Thr Lys Ile
545 550 555 560

Asp Asp Lys Gly Val Ser Phe Val Asp Ser Ser Gly Gln Ala Lys Ala
565 570 575

Asn Thr Pro Val Leu Ser Ala Asn Gly Leu Asp Leu Gly Gly Lys Val
580 585 590

Ile Ser Asn Val Gly Lys Gly Thr Lys Asp Thr Asp Ala Ala Asn Val
595 600 605

Gln Gln Leu Asn Glu Val Arg Asn Leu Leu Gly Leu Gly Asn Ala Gly
610 615 620

Asn Asp Asn Ala Asp Gly Asn Gln Val Asn Ile Ala Asp Ile Lys Lys
625 630 635 640

Asp Pro Asn Ser Gly Ser Ser Ser Asn Arg Thr Val Ile Lys Ala Gly
645 650 655

Thr Val Leu Gly Gly Lys Gly Asn Asn Asp Thr Glu Lys Leu Ala Thr
660 665 670

Gly Gly Ile Gln Val Gly Val Asp Lys Asp Gly Asn Ala Asn Gly Asp
675 680 685

Leu Ser Asn Val Trp Val Lys Thr Gln Lys Asp Gly Ser Lys Lys Ala
690 695 700

Leu Leu Ala Thr Tyr Asn Ala Ala Gly Gln Thr Asn Tyr Leu Thr Asn
705 710 715 720

Asn Pro Ala Glu Ala Ile Asp Arg Ile Asn Glu Gln Gly Ile Arg Phe
725 730 735

Substitute SeqListing.txt

Phe His Val Asn Asp Gly Asn Gln Glu Pro Val Val Gln Gly Arg Asn
740 745 750

Gly Ile Asp Ser Ser Ala Ser Gly Lys His Ser Val Ala Ile Gly Phe
755 760 765

Gln Ala Lys Ala Asp Gly Glu Ala Ala Val Ala Ile Gly Arg Gln Thr
770 775 780

Gln Ala Gly Asn Gln Ser Ile Ala Ile Gly Asp Asn Ala Gln Ala Thr
785 790 795 800

Gly Asp Gln Ser Ile Ala Ile Gly Thr Gly Asn Val Val Ala Gly Lys
805 810 815

<210> 49
<211> 720
<212> DNA
<213> Moraxella catarrhalis

<400> 49
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<210> 50
<211> 60
<212> PRT
<213> Moraxella catarrhalis

<400> 50
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20 25 30

Cys Ala Thr Gly Gln Val Gly Ser Val Cys Thr Leu Ser Phe Ala Arg
35 40 45

Ile Ala Ala Leu Ala Val Leu Val Ile Gly Ala Thr
50 55 60

<210> 51
<211> 5
<212> PRT
<213> Moraxella cattarhalis

<400> 51
Val Val Ala Gly Lys

Substitute SeqListing.txt

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5